

DNA sequence 489 b.p. atgagaatttcg ... aacactctctga linear

1/1 31/11  
 atg aga att tgg aaa cca cat tgg aga agt att tcc atc cag tgc tac tgg tgt tta ctt  
 Met arg ile ser lys pro his leu arg ser ile ser ile gln cys tyr leu cys leu leu  
 61/21 91/31  
 cta aac agt cat ttt cta act gaa gct ggc att cat gtc ttc att ttg ggc tgt ttc agt  
 leu asn ser his phe leu thr glu ala gly ile his val phe ile leu gly cys phe ser  
 121/41 151/51  
 gca ggg ctt cct aaa aca gaa gcc aac tgg gtg aat gta ata agt gat ttg aaa aaa att  
 ala gly leu pro lys thr glu ala asn trp val asn val ile ser asp leu lys lys ile  
 181/61 211/71  
 gaa gat ctt att caa tct atg cat att gat gct act tta tat acg gaa agt gat gtc cac  
 glu asp leu ile gln ser met his ile asp ala thr leu tyr thr glu ser asp val his  
 241/81 271/91  
 ccc agt tgt aaa gta aca gca atg aag tgc ttt ctc ttg gag tta caa gtt att tca ctt  
 pro ser cys lys val thr ala met lys cys phe leu leu glu leu gln val ile ser leu  
 301/101 331/111  
 gag tcc gga gat gca agt att cat gat aca gta gaa aat ctg atc atc cta gca aac aac  
 glu ser gly asp ala ser ile his asp thr val glu asn leu ile ile leu ala asn asn  
 361/121 391/131  
 agt ctg tct tct aat ggg aat gca aca gaa tct gga tgc aaa gaa tgt gag gaa ctg gag  
 ser leu ser ser asn gly asn val thr glu ser gly cys lys glu cys glu glu leu glu  
 421/141 451/151  
 gaa aaa aat att aaa gaa ttt ttg gac agt ttt gta cat att gtc gac atg ttc atc aac  
 glu lys asn ile lys glu phe leu asp ser phe val his ile val asp met phe ile asn  
 481/161  
 att tct tga (SEQ ID NO:1)  
 thr ser opa (SEQ ID NO:2)

Fig. 1

DNA sequence 489 b.p. atgagaattctg ... aacacttcttga linear

1/1  
 atg aga att tgg aaa cca cat ttg aga agt att tcc atc cag tgc tac ttg tgt tta ctt  
 Met arg ile ser lys pro his leu arg ser ile ser ile gln cys tyr leu cys leu leu  
 31/11  
 61/21  
 cta aac agt cat ttt cta act gaa gct ggc att cat gtc ttc att ttg ggc tgt ttc agt  
 leu asn ser his phe leu thr glu ala gly ile his val phe ile leu gly cys phe ser  
 91/31  
 121/41  
 gca ggg ctt cct aaa aca gaa gcc aac tgg gtg aat gta ata agt gat ttg aaa aaa att  
 ala gly leu pro lys thr glu ala asn trp val asn val ile ser asp leu lys lys ile  
 151/51  
 181/61  
 gaa gat ctt att caa tct atg cat att gat gct act tta tat acg gaa agt gat gtt cac  
 glu asp leu ile gln ser met his ile asp ala thr leu tyr thr glu ser asp val his  
 211/71  
 241/81  
 ccc agt tgc aaa gta aca gca atg aag tgc ttt ctc ttg gag tta caa gtt att tca ctt  
 pro ser cys lys val thr ala met lys cys phe leu leu glu leu gln val ile ser leu  
 271/91  
 301/101  
 gag tcc gga gat gca agt att cat gat aca gta gaa aat ctg atc atc cta gca aac aac  
 glu ser gly asp ala ser ile his asp thr val glu asn leu ile ile leu ala asn asn  
 331/111  
 361/121  
 agt ttg tct tct aat ggg aat gta aca gaa tct gga tgc aaa gaa tgt gag gaa ctg gag  
 ser leu ser ser asn gly asn val thr glu ser gly cys lys glu cys glu glu leu glu  
 391/131  
 421/141  
 gaa aaa aat att aaa gaa ttt ttg cag agt ttt gta cat att gtc caa atg ttc atc aac  
 glu lys asn ile lys glu phe leu gln ser phe val his ile val gln met phe ile asn  
 451/151  
 481/161  
 act tct tga (SEQ ID NO: 3)  
 thr ser opa (SEQ ID NO: 4)

Fig. 2

0955313-051401

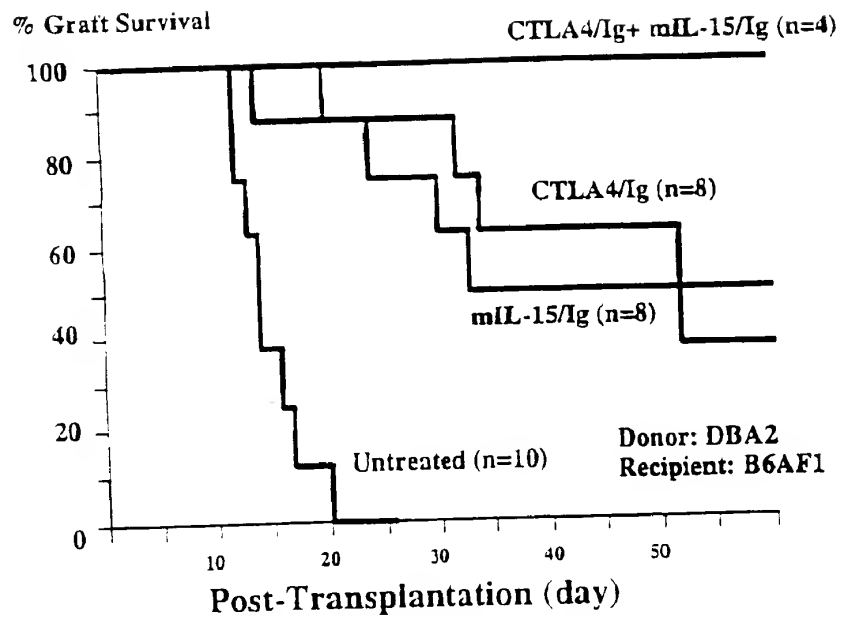


Fig. 3